Remarks

The amendments presented herein are being made to conform the application with the provisions of 37 CFR §§1.821 through 1.825 and to correct inadvertent typographical errors throughout the specification. Support for the amendment to Figure 1 can be found, for example, at page 2, lines 24-29, and page 3, lines 1-2. Support for new SEQ ID NOs. 10-16 can be found in the references and accession numbers cited at page 5, lines 3-5, of the subject specification. The cited references and accession numbers disclose the sequences shown in SEQ ID NOs. 10-16. Applicants note that the subject application incorporates by reference all publications cited therein (see page 12, lines 28-30, of the subject specification). I hereby certify that no new material is being added by this submission.

The Commissioner is hereby authorized to charge any fees which may be required to Deposit Account No. 19-0065.

Respectfully submitted,

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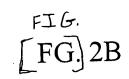
Gainesville, FL 32606-6669

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Attachments: New pages 1-47 (Sequence Listing) of the subject specification; new Figures 1 and

2B.

catcgataaa .	taaaaactg	c agcccg	gggg ato	ccatagt	tg taaaa	aattc at	a atg Met 1	1676
gaa gtc tgc Glu Val Cys	aat tgt Asn Cys 5	att gaa Ile Glu	ccg caa Pro Gln 10	tgg co	ca gcg g ro Ala <i>P</i>	gat gaa t Asp Glu I 15	tg tta Leu Leu	1724
atg aaa tac Met Lys Tyr 20	caa tac Gln Tyr	atc tcc Ile Ser	gat ttc Asp Phe 25	ttc a	TC 1770 -	att gcg t Ile Ala T 30	at ttt Tyr Phe	1772
tcg att cct Ser Ile Pro 35	ctt gag Leu Glu	ttg att Leu Ile 40	tac ttt Tyr Phe	gtg a Val L	ag aaa t ys Lys : 45	tca gcc q Ser Ala \	gtg ttt Val Phe	1820
ccg tat aga Pro Tyr Arg 50	tgg gta Trp Val	ctt gtt Leu Val 55	cag ttt Gln Phe	: GIY F	oct ttt Ala Phe 50	atc gtt o Ile Val I	ctt tat Leu Tyr 65	1868
gga gca act Gly Ala Thi	cat ctt His Leu 70	att aac Ile Asn	tta tgo Leu Tr	g act to Thr E 75	ttc act Phe Thr	acg cat Thr His	tcg aga Ser Arg 80	1916
acc gtg gc Thr Val Al	g ctt gtg a Leu Val 85	atg act Met Thr	acc gco Thr Ala	g aag q a Lys \	gtg tta Val Leu	acc gct Thr Ala 95	gtt gtc Val Val	1964
tcg tgt gc Ser Cys Al	a Thr Ala	g ttg atg A Leu Met	ctt gt Leu Va 105	t cat a	att att Ile Ile	cct gat Pro Asp 110	ctt ttg Leu Leu	2012
agt gtt aa Ser Val Ly 115	g act cgo s Thr Aro	g gag ctt g Glu Lei 120	i Phe Le	g aaa u Lys .	aat aaa Asn Lys 125	gct gct Ala Ala	gag ctc Glu Leu	2060
gat aga ga Asp Arg Gl 130	a atg gga u Met Gly	a ttg att y Leu Ile 135	arg in	I GIII	gaa gaa Glu Glu 140	acc gga Thr Gly	agg cat Arg His 145	2108
gtg aga at Val Arg Me	g ttg ac t Leu Th 15	r His GI	g att ag u Ile Ar	a agc g Ser 155	act tta Thr Leu	gat aga Asp Arg	cat act His Thr 160	2156
att tta aa Ile Leu L	ng act ac ys Thr Th 165	a ctt gt r Leu Va	t gag ct l Glu Le 1	su Oly	agg aca Arg Thr	tta gct Leu Ala 175	ttg gag Leu Glu	2204
gag tgt g Glu Cys A 1	ca ttg tg la Leu Tr 30	g atg cc p Met Pr	t act act act of Thr A	ga act rg Thr	ggg tta Gly Leu	gag cta Glu Leu 190	cag ctt Gln Leu	2252





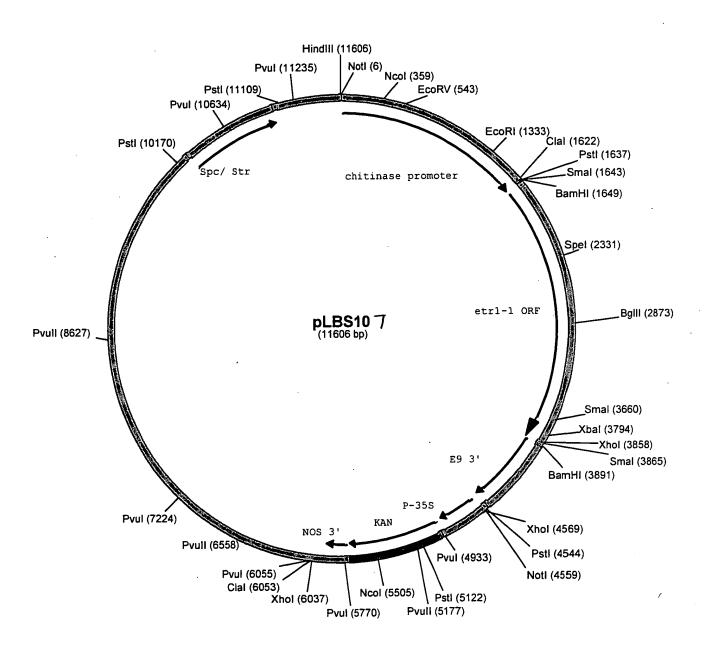


FIG. 1